

-NKEQHLSSENPVDTNSDNNLFTDTDLK-	503-529	<u>18</u>
-EMDDVAIEDEVLEQLFK-	552-558	<u>19</u>
-MDIETNDTFSDEAVPESSK-	595-613	<u>20</u>
-ELKEDSWAK-	625-635	<u>21</u>
-KLLLTEFR-	653-660	<u>22</u>
-NPSGINDDYGQLK- ^c	671-683	<u>23</u>
-EESLADDLFR-	736-745	<u>24</u>

^aMass spectrometry is unable to distinguish L/I and Q/K. The indication of L, I, Q, and K residues in the peptide sequences is based on the p95 ORF.

^bNumbering system is based on the ORF of the p95 cDNA.

^cThis peptide was sequenced by tandem mass spectrometry and used for EST database searches. The remaining peptides corresponded to the derived amino acid sequence from the p95 ORF.
